OIPE

#11

RAW SEQUENCE LISTING DATE: 08/15/2001 PATENT APPLICATION: US/09/559,021 TIME: 07:33:25

Input Set : A:\Seq Listing.txt

Output Set: N:\CRF3\08132001\I559021.raw

## ENTERED

```
3 <110> APPLICANT: WOLFF, JON
             SOKOLOFF, ALEXANDER
      6 <120> TITLE OF INVENTION: PROCESS FOR UTILIZING EPITOPES RECOGNIZED BY NATURAL
ANTIBODIES
     8 <130> FILE REFERENCE: MIRUS.014.02
    10 <140> CURRENT APPLICATION NUMBER: 09/559021
    11 <141> CURRENT FILING DATE: 2000-04-27
    13 <160> NUMBER OF SEQ ID NOS: 120
    15 <170> SOFTWARE: PatentIn version 3.1
    17 <210> SEQ ID NO: 1
    18 <211> LENGTH: 11
    19 <212> TYPE: PRT
    20 <213> ORGANISM: Bacteriophage T7
    22 <400> SEQUENCE: 1
    24 Phe Gln Ser Gly Val Met Leu Gly Asp Pro Asn
     28 <210> SEQ ID NO: 2
    29 <211> LENGTH: 27
    30 <212> TYPE: PRT
    31 <213> ORGANISM: Bacteriophage T7
    33 <400> SEQUENCE: 2
    35 Phe Gln Ser Gly Val Met Leu Gly Asp Pro Asn Ser Asp Gly Ala Leu
    39 Arg Gln Ser Gly Arg Gly Lys Ser Ser Arg Pro
                    20
     43 <210> SEQ ID NO: 3
     44 <211> LENGTH: 23
     45 <212> TYPE: PRT
     46 <213> ORGANISM: Bacteriophage T7
     48 <400> SEQUENCE: 3
    50 Phe Gln Ser Gly Val Met Leu Gly Asp Pro Asn Ser Ser Val Asp
    51 1
                                            10
    54 Lys Leu Ala Ala Leu Glu
                    20
    58 <210> SEQ ID NO: 4
     59 <211> LENGTH: 8
     60 <212> TYPE: PRT
     61 <213> ORGANISM: Bacteriophage T7
     63 <400> SEQUENCE: 4
     65 Ala Ala Gly Ala Val Val Phe Gln
     69 <210> SEQ ID NO: 5
     70 <211> LENGTH: 343
     71 <212> TYPE: PRT
     72 <213> ORGANISM: Bacteriophage T7
    74 <400> SEQUENCE: 5
     76 Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Thr Asn Gln Gly Lys
     77 1
```

Input Set : A:\Seq Listing.txt

```
80 Gly Val Val Ala Ala Gly Asp Lys Leu Ala Leu Phe Leu Lys Val Phe
84 Gly Gly Glu Val Leu Thr Ala Phe Ala Arg Thr Ser Val Thr Thr Ser
                               40
88 Arg His Met Val Arg Ser Ile Ser Ser Gly Lys Ser Ala Gln Phe Pro
                           55
92 Val Leu Gly Arg Thr Gln Ala Ala Tyr Leu Ala Pro Gly Glu Asn Leu
                       70
                                           75
96 Asp Asp Lys Arg Lys Asp Ile Lys His Thr Glu Lys Val Ile Thr Ile
100 Asp Gly Leu Leu Thr Ala Asp Val Leu Ile Tyr Asp Ile Glu Asp Ala
101
                100
104 Met Asn His Tyr Asp Val Arg Ser Glu Tyr Thr Ser Gln Leu Gly Glu
                                120
108 Ser Leu Ala Met Ala Ala Asp Gly Ala Val Leu Ala Glu Ile Ala Gly
                            135
112 Leu Cys Asn Val Glu Ser Lys Tyr Asn Glu Asn Ile Glu Gly Leu Gly
                                            155
                        150
116 Thr Ala Thr Val Ile Glu Thr Thr Gln Asn Lys Ala Ala Leu Thr Asp
                                        170
                    165
120 Gln Val Ala Leu Gly Lys Glu Ile Ile Ala Ala Leu Thr Lys Ala Arg
                                    185
124 Ala Ala Leu Thr Lys Asn Tyr Val Pro Ala Ala Asp Arg Val Phe Tyr
                                200
                                                    205
            195
128 Cys Asp Pro Asp Ser Tyr Ser Ala Ile Leu Ala Ala Leu Met Pro Asn
                            215
132 Ala Ala Asn Tyr Ala Ala Leu Ile Asp Pro Glu Lys Gly Ser Ile Arg
                                            235
                       230
136 Asn Val Met Gly Phe Glu Val Val Glu Val Pro His Leu Thr Ala Gly
                                        250
                   245
140 Gly Ala Gly Thr Ala Arg Glu Gly Thr Thr Gly Gln Lys His Val Phe
                                    265
141
               260
144 Pro Ala Asn Lys Gly Glu Gly Asn Val Lys Val Ala Lys Asp Asn Val
           275
                                280
148 Ile Gly Leu Phe Met His Arg Ser Ala Val Gly Thr Val Lys Leu Arg
                            295
152 Asp Leu Ala Leu Glu Arg Ala Arg Arg Ala Asn Phe Gln Ala Asp Gln
                        310
                                            315
156 Ile Ile Ala Lys Tyr Ala Met Gly His Gly Gly Leu Arg Pro Glu Ala
                    325
157
160 Ala Gly Ala Val Val Phe Gln
                340
164 <210> SEQ ID NO: 6
165 <211> LENGTH: 9
166 <212> TYPE: PRT
167 <213> ORGANISM: Bacteriophage T7
169 <400> SEQUENCE: 6
171 Ala Ala Gly Ala Val Val Phe Gln Ser
172 1
```

Input Set : A:\Seq Listing.txt

```
175 <210> SEO ID NO: 7
176 <211> LENGTH: 10
177 <212> TYPE: PRT
178 <213> ORGANISM: Bacteriophage T7
180 <400> SEQUENCE: 7
182 Ala Ala Gly Ala Val Val Phe Ser Gln Val
183 1
                    5
186 <210> SEQ ID NO: 8
187 <211> LENGTH: 9
188 <212> TYPE: PRT
189 <213> ORGANISM: Bacteriophage T7
191 <400> SEQUENCE: 8
193 Glu Ala Ala Gly Ala Val Val Phe Gln
194 1
                    5
197 <210> SEQ ID NO: 9
198 <211> LENGTH: 13
199 <212> TYPE: PRT
200 <213> ORGANISM: phage SV40
202 <400> SEQUENCE: 9
204 Cys Gly Tyr Gly Pro Lys Lys Arg Lys Val Gly Gly
205 1
208 <210> SEQ ID NO: 10
209 <211> LENGTH: 39
210 <212> TYPE: PRT
211 <213> ORGANISM: phage SV40
213 <400> SEQUENCE: 10
215 Cys Lys Lys Ser Ser Ser Asp Asp Glu Ala Thr Ala Asp Ser Gln
                                        10
219 His Ser Thr Pro Pro Lys Lys Lys Arg Lys Val Glu Asp Pro Lys Asp
220
                                    25
                20
223 Phe Pro Ser Glu Leu Leu Ser
224
            35
227 <210> SEQ ID NO: 11
228 <211> LENGTH: 38
229 <212> TYPE: PRT
230 <213> ORGANISM: phage SV40
232 <400> SEQUENCE: 11
234 Cys Lys Lys Trp Asp Asp Glu Ala Thr Ala Asp Ser Gln His
235 1
238 Ser Thr Pro Pro Lys Lys Lys Arg Lys Val Glu Asp Pro Lys Asp Phe
                                    25
239
                20
242 Pro Ser Glu Leu Leu Ser
243
            35
246 <210> SEQ ID NO: 12
247 <211> LENGTH: 32
248 <212> TYPE: PRT
249 <213> ORGANISM: M9 Protein
251 <400> SEQUENCE: 12
253 Cys Tyr Asn Asp Phe Gly Asn Tyr Asn Asn Gln Ser Ser Asn Phe Gly
```

Input Set : A:\Seq Listing.txt

```
254 1
                                         10
                                                             15
257 Pro Met Lys Gln Gly Gly Asn Phe Gly Gly Arg Ser Ser Gly Pro Tyr
                20
                                    25
261 <210> SEQ ID NO: 13
262 <211> LENGTH: 10
263 <212> TYPE: PRT
264 <213> ORGANISM: E1A Adenovirus
266 <400> SEQUENCE: 13
268 Cys Lys Arg Gly Pro Lys Arg Pro Arg Pro
269 1
272 <210> SEQ ID NO: 14
273 <211> LENGTH: 22
274 <212> TYPE: PRT
275 <213> ORGANISM: Nucleoplasmin
277 <400> SEQUENCE: 14
279 Cys Lys Lys Ala Val Lys Arg Pro Ala Ala Thr Lys Lys Ala Gly Gln
                                         10
280 1
283 Ala Lys Lys Lys Leu
284
                20
287 <210> SEQ ID NO: 15
288 <211> LENGTH: 14
289 <212> TYPE: PRT
290 <213> ORGANISM: c-myc
292 <400> SEQUENCE: 15
294 Cys Lys Lys Gly Pro Ala Ala Lys Arg Val Lys Leu Asp
295 1
298 <210> SEQ ID NO: 16
299 <211> LENGTH: 4
300 <212> TYPE: PRT
301 <213> ORGANISM: Bacteriophage T7
303 <400> SEQUENCE: 16
305 Phe Ser Gln Val
306 1
309 <210> SEQ ID NO: 17
310 <211> LENGTH: 4
311 <212> TYPE: PRT
312 <213> ORGANISM: endoplasmic reticulum proteins
314 <400> SEQUENCE: 17
316 Lys Asp Glu Leu
317 1
320 <210> SEQ ID NO: 18
321 <211> LENGTH: 4
322 <212> TYPE: PRT
323 <213> ORGANISM: Bacteriophage T7
325 <400> SEQUENCE: 18
327 Gln Val Thr Lys
328 1
331 <210> SEO ID NO: 19
332 <211> LENGTH: 8
```

Input Set : A:\Seq Listing.txt

```
333 <212> TYPE: PRT
334 <213> ORGANISM: Bacteriophage T7
336 <400> SEQUENCE: 19
338 Val Val Val Glu Ser Val Pro Lys
339 1
342 <210> SEQ ID NO: 20
343 <211> LENGTH: 6
344 <212> TYPE: PRT
345 <213> ORGANISM: Bacteriophage T7
347 <400> SEQUENCE: 20
349 Ala Arg Pro Val Gln Lys
350 1
353 <210> SEQ ID NO: 21
354 <211> LENGTH: 8
355 <212> TYPE: PRT
356 <213> ORGANISM: Bacteriophage T7
358 <400> SEQUENCE: 21
360 Gln Leu Val Arg Val Ile Ser Arg
361 1
364 <210> SEQ ID NO: 22
365 <211> LENGTH: 4
366 <212> TYPE: PRT
367 <213> ORGANISM: Bacteriophage T7
369 <400> SEQUENCE: 22
371 Gly Arg Leu Lys
372 1
375 <210> SEQ ID NO: 23
376 <211> LENGTH: 5
377 <212> TYPE: PRT
378 <213> ORGANISM: Bacteriophage T7
380 <400> SEQUENCE: 23
382 Ala Phe Thr Asn Lys
383 1
386 <210> SEQ ID NO: 24
387 <211> LENGTH: 6
388 <212> TYPE: PRT
389 <213> ORGANISM: Bacteriophage T7
391 <400> SEQUENCE: 24
393 Val Thr Pro Gln Val Lys
394 1
397 <210> SEQ ID NO: 25
398 <211> LENGTH: 7
399 <212> TYPE: PRT
400 <213> ORGANISM: Bacteriophage T7
402 <400> SEQUENCE: 25
404 Asp Asn Thr Pro Lys Thr Lys
405 1
408 <210> SEQ ID NO: 26
409 <211> LENGTH: 12
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/559,021

DATE: 08/15/2001 TIME: 07:33:26

Input Set : A:\Seq Listing.txt
Output Set: N:\CRF3\08132001\I559021.raw